## Grant Anderson LLP

RECEIVED CENTRAL FAX CENTER MAY 0 6 2008

6540 LUSK BLVD., SUITE C210 SAN DIEGO, CALIFORNIA 92121 TOBEY M. TAM
Patent Agent
Direct (858) 248-0178
Facsimile (858) 623-3224
Email ttam@granllp.com

## **FACSIMILE**

Re:	US App. No. 10/723,681 – proposed claim amendments		
Telephone No.	571-272-0752	Pages:	8 (including cover page)
Facsimile No.	571-273-8300	Date:	May 6, 2008
То:	Examiner Jehanne Sitton	From:	Tobey M. Tam

Examiner Sitton,

Please find the following 7 pages a proposed claim amendment for US App. No. 10/723,681. We look forward to our telephone interview Thurs, May 8, 2008 at 11am EST.

If you have any questions beforehand, do not hesitate to contact me.

Thank you, Tobey Tam

## RECEIVED CENTRAL FAX CENTER

MAY 0 6 2008

Confidential - US App No. 10/723,681 For Discussion Purposes Only Not for Filing

Interview with the Examiner Sitton, Bruce Grant and Tobey Tam on Thursday, May 8, 2008 at 11am EST.

Proposed claim amendments for US Application No. 10/723,681 (our reference SEQ-4069-UT):

1 (currently amended). A method for identifying a subject at risk of breast cancer, which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a human subject, wherein one or more polymorphic variations are detected in one or more regions selected from a region between about at one or more chromosome positions selected from the group consisting of chromosome positions 87330326, 87332557, 87332861, 87333099, 87333312, 87333569, 87341627, 87341722, and 87342924 87330326 to about chromosome position 87342924, a region between about chromosome position 87352676 to about chromosome position 87369072, a region between about chromosome position 87311012 to about chromosome position 87314967 and a region between about chromosome position 87320855, wherein each chromosome position is according to Build 33 of the GenBank database human genome sequence

whereby the presence of the one or more polymorphic variations is indicative of the subject is identified as being at risk of breast cancer based on the presence or absence of the one or more polymorphic variations associated with breast cancer.

2 (original). The method of claim 1, which further comprises obtaining the nucleic acid sample from the subject.

3-18 (cancelled).